

Attorney Docket No.: UIC0005US.NP
Inventors: Kumar et al.
Serial No.: Not yet assigned
Filing Date: Herewith
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Amendments to the Drawings:

The attached sheet of drawings includes changes to Figure 3D. This sheet, which includes Figures 3D, previously listed SEQ ID No. 81 in duplicate.

Attachment: Replacement Sheet
Annotated Sheet Showing Changes

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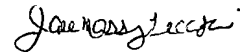
Remarks/Arguments

Claims 1-11 are pending in the instant application. Claims 1-11 have been canceled by this preliminary amendment. New claims 12-20 have been added. Support for the newly presented claims is found throughout the specification as filed. An amended sequence listing and replacement Figure 3D are also provided. SEQ ID NO:159 is referred to in the specification at page 5, line 8 as Genbank Accession Number: M85079. A SEQ ID NO. has been assigned to facilitate search and examination. SEQ ID NO:160 is a designated coding sequence located within SEQ ID NO:159. PatentIn 3.3 automatically generates a corresponding amino acid SEQ ID NO. for SEQ ID NO:159. Two different sequences were identified as SEQ ID NO:81 in the specification as filed. The Sequence Listing and Figure 3D have been amended to correct this inadvertent error. In addition, Applicants have amended the specification to include the claim of priority and to include the acknowledgment of

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U.S. Government rights in this invention. No new matter is added by this amendment. Entry of this amendment is respectfully requested.

Respectfully submitted,



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Date: **February 10, 2006**

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Figure 3D

Target sequence 23: AACATCAACCACAACACAGAG (SEQ ID NO: 78)

Position in gene sequence: 1035

GC content: 42.9%

Sense strand siRNA: CAUCAACCACAACACAGAGtt (SEQ ID NO: 79)

Antisense strand siRNA: CUCUGUGUUGUGGUUGAUGtt (SEQ ID NO: 80)

Target sequence 24: AAGCTGAAGCAGAACACTTCA (SEQ ID NO: 81)

Position in gene sequence: 1119

GC content: 42.9%

Sense strand siRNA: GCUGAAGCAGAACACUUCAtt (SEQ ID NO: 81)

Antisense strand siRNA: UGAAGUGUUCUGCUUCAGCtt (SEQ ID NO: 82)

Target sequence 25: AAGCAGAACACTTCAGAGCAG (SEQ ID NO: 83)

Position in gene sequence: 1125

GC content: 47.6%

Sense strand siRNA: GCAGAACACUUCAGAGCAGtt (SEQ ID NO: 84)

Antisense strand siRNA: CUGCUCUGAAGUGUUCUGCtt (SEQ ID NO: 85)

Target sequence 26: AACACTTCAGAGCAGTTGAG (SEQ ID NO: 86)

Position in gene sequence: 1131

GC content: 42.9%

Sense strand siRNA: CACUUCAGAGCAGUUUGAGtt (SEQ ID NO: 87)

Antisense strand siRNA: CUCAAACUGCUCUGAAGUGtt (SEQ ID NO: 88)

Target sequence 27: AAGATCTTCCCTATGAGGAG (SEQ ID NO: 89)

Position in gene sequence: 1164

GC content: 42.9%

Sense strand siRNA: GAUCUUUCCCUAUGAGGAGtt (SEQ ID NO: 90)

Antisense strand siRNA: CUCCUCAUAGGGAAAGAUCtt (SEQ ID NO: 91)

Target sequence 28: AAGACAGAGAAGGACATCTTC (SEQ ID NO: 92)

Position in gene sequence: 1197

GC content: 42.9%

Sense strand siRNA: GACAGAGAAGGACAUCUUCtt (SEQ ID NO: 93)

Antisense strand siRNA: GAAGAUGUCCUUCUCUGUCtt (SEQ ID NO: 94)

Target sequence 29: AAGGACATCTTCTCAGACATC (SEQ ID NO: 95)

Position in gene sequence: 1206

GC content: 42.9%

Sense strand siRNA: GGACAUCUUCUCAGACAUCtt (SEQ ID NO: 96)

Antisense strand siRNA: GAUGUCUGAGAAGAUGUCtt (SEQ ID NO: 97)

Correcting
duplicate
SEQ ID NO: 81
to
SEQ ID NO:
161